

SEQUENCE LISTING

<110> TRANSGENE SA et al.

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<170> PatentIn Ver. 2.1

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Val Leu Thr Val Val Thr Gly Ser Gly His Ala Ser Ser Thr Pro Gly
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Gly Glu Lys Glu Thr Ser Ala Thr Gln Arg Ser Ser Val Pro Ser Ser
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Thr Glu Lys Asn Ala Val Ser Met Thr Ser Ser Val Leu Ser Ser His
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Ser Pro Gly Ser Gly Ser Ser Thr Thr Gln Gly Gln Asp Val Thr Leu
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Ala Pro Ala Thr Glu Pro Ala Ser Gly Ser Ala Ala Thr Trp Gly Gln
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Asp Val Thr Ser Val Pro Val Thr Arg Pro Ala Leu Gly Ser Thr Thr
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Pro Pro Ala His Asp Val Thr Ser Ala Pro Asp Asn Lys Pro Ala Pro
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Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr
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gcc ccg gac acc agg ccg gcc ccg ggc tcc acc gcc ccc ccc ggc gac Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His 165	170	175		585
ggg gtc acc tcg gcc ccg gac aac agg ccg gcc ttg ggc tcc act gcc Gly Val Thr Ser Ala Pro Asp Asn Arg Pro Ala Leu Gly Ser Thr Ala 180	185	190		633
cct cca gtc cac aat gtc acc tcg gcc tca ggc tct gca tca ggc tca Pro Pro Val His Asn Val Thr Ser Ala Ser Gly Ser Ala Ser Gly Ser 195	200	205		681
gtc tct act ctg gtg cac aac ggc acc tct gcc agg gct acc aca acc Ala Ser Thr Leu Val His Asn Gly Thr Ser Ala Arg Ala Thr Thr Thr 210	215	220		729
cca gcc agc aag agc act cca ccc agc att ccc agc gag cac cac tct gat Pro Ala Ser Lys Ser Thr Pro Pro Ser Ile Pro Ser His His Ser Asp 225	230	235	240	777
act cct acc acc ctt gcc agc cat agc acc aag act gat gcc agt agc Thr Pro Thr Thr Leu Ala Ser His Ser Thr Lys Thr Asp Ala Ser Ser 245	250	255		825
act cac cat agc acg gta cct cct ctc acc tcc tcc aat cac agc act Thr His His Ser Thr Val Pro Pro Leu Thr Ser Ser Asn His Ser Thr 260	265	270		873
tct ccc cag ttg tct act ggg gtc tct ttc ttc ctg tct ttt cac Ser Pro Gln Leu Ser Thr Gly Val Ser Phe Phe Phe Leu Ser Phe His 275	280	285		921
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tat aaa caa ggg ggt ttt ctg ggc ctc tcc aat att aag ttc agg cca Tyr Lys Gln Gly Gly Phe Leu Gly Leu Ser Asn Ile Lys Phe Arg Pro 325	330	335		1065
gga tct gtg gtg gta caa ttg act ctg gcc ttc cga gaa ggt acc atc Gly Ser Val Val Gln Leu Thr Leu Ala Phe Arg Glu Gly Thr Ile 340	345	350		1113
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gcc tct ega tat aac ctg acg atc tca gac gtc agc gtg agt cat gtg Ala Ser Arg Tyr Asn Leu Thr Ile Ser Asp Val Ser Val Ser His Val 370	375	380		1209
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 Ile Ala Leu Leu Val Leu Val Cys Val Leu Val Ala Leu Ala Ile Val 1305
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 Tyr Leu Ile Ala Leu Ala Val Cys Gln Cys Arg Arg Lys Asn Tyr Gly 1353
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cag ctg gac atc ttt cca gcc cgg gat acc tac cat cct atc agc gag
 Gln Leu Asp Ile Phe Pro Ala Arg Asp Thr Tyr His Pro Met Ser Glu 1401
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 Tyr Pro Thr His Thr His Gly Arg Tyr Val Pro Ser Ser Thr 1449
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 Asp Arg Ser Pro Tyr Glu Lys Val Ser Ala Gly Asn Gly Ser Ser 1497
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Thr Glu Lys Asn Ala Val Ser Met Thr Ser Ser Val Leu Ser Ser His
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Ser Pro Gly Ser Gly Ser Ser Thr Thr Gln Gly Gln Asp Val Thr Leu
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Ala Pro Ala Thr Glu Pro Ala Ser Gly Ser Ala Ala Thr Trp Gly Gln
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Pro Pro Ala His Asp Val Thr Ser Ala Pro Asp Asn Lys Pro Ala Pro
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Arg Pro Pro Pro Gly Ser Thr Ala Pro Ala Ala His Gly Val Thr Ser
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Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His
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 Pro Pro Val His Asn Val Thr Ser Ala Ser Gly Ser Ala Ser Gly Ser
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 Ala Ser Thr Leu Val His Asn Gly Thr Ser Ala Arg Ala Thr Thr Thr
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 Pro Ala Ser Lys Ser Thr Pro Pro Ser Ile Pro Ser His His Ser Asp
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 Thr Pro Thr Thr Leu Ala Ser His Ser Thr Lys Thr Asp Ala Ser Ser
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 Thr His His Ser Thr Val Pro Pro Leu Thr Ser Ser Asn His Ser Thr
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 Ile Ser Asn Leu Gln Phe Asn Ser Ser Leu Glu Asp Pro Ser Thr Asp
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 Tyr Tyr Gln Glu Leu Gln Arg Asp Ile Ser Gln Met Phe Leu Gln Ile
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 Tyr Lys Gln Gly Gly Phe Leu Gly Leu Ser Asn Ile Lys Phe Arg Pro
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Pro Pro Ala His Asp Val Thr Ser Ala Pro Asp Asn Lys Pro Ala Pro
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Val His Asn Gly Thr Ser Ala Arg Ala Thr Thr Pro Ala Ser Lys

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Leu Ala Ser His Ser Thr Lys Thr Asp Ala Ser Ser Thr His His Ser
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Tyr Glu Lys Val Ser Ala Gly Asn Gly Gly Ser Ser Leu Ser Tyr Thr
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